

*Kidney International, Vol. 57 (2000), pp. 1895–1904*

# Exposure of endothelial cells to recombinant human erythropoietin induces nitric oxide synthase activity

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## Exposure of endothelial cells to recombinant human erythropoietin induces nitric oxide synthase activity.

**Background.** Anemic patients with chronic renal failure receiving recombinant human erythropoietin (rHuEPO) therapy frequently develop hypertension through an unknown mechanism. We hypothesize that EPO receptors (EPORs) on endothelial cells (ECs) in various sites of vasculature may mediate the activities of nitric oxide synthase (NOS) and/or the release of endothelin-1 (ET-1), contributing to blood pressure changes. We tested this hypothesis using primary cultures of ECs obtained from human coronary artery (HCAEC), pulmonary artery (HPAEC), dermis (HDEC), and umbilical vein (HUVEC).

**Methods.** EPORs were measured by  $^{125}\text{I}$ -EPO binding. The effect of EPO on EPOR, ET-1, and NOS mRNA levels was assessed by quantitative reverse transcription-polymerase chain reaction. Cellular NOS activity and ET-1 release into the medium was measured by the NOSdetect assay and by radioimmunoassay kits.

**Results.** Short-term (4 h) treatment with EPO (4 U/mL) did not change the number or affinity of EPOR per cell. Neither were there any changes in the amount of EPOR, ET-1, and NOS transcripts (cDNA/ $\mu\text{g}$  of mRNA) nor in ET-1 release and NOS activity. In HUVEC only, 24-hour exposure to EPO caused a threefold increase in NOS transcript. In other cells, EPO treatment for six days increased NOS activity by twofold to fourfold.

**Conclusions.** We show that upon extended exposure, EPO induces NOS activity but does not affect ET-1 release. These findings indicate that the hypertensive effect of EPO is not likely to be caused by a direct effect on ECs.

Erythropoietin (EPO) is a glycoprotein hormone that is the primary regulator of erythropoiesis [1]. EPO is produced by the kidney in adults and by the liver in fetal life [1, 2]. EPO functions through its interaction with a single chain cell surface receptor of the cytokine receptor superfamily [3, 4]. The EPO receptor (EPOR) on ery-

throid progenitor cells is the primary target for EPO binding [5, 6].

In hematopoietic cells, EPOR mRNA is expressed at moderate levels [7]. EPOR or EPOR mRNA are also expressed on nonhematopoietic cells, including human umbilical vein endothelial cell (HUVEC) [8, 9], rat brain capillary endothelial cells (ECs) [10], murine hippocampal and cerebrotectal areas [11], and primary cultured hippocampal and cortical neurons [12, 13]. The EPORs are functional in HUVEC and mouse brain.

The angiogenic effect of EPO has been studied in ovariectomized mice in which an injection of EPO into the uterine cavity promoted blood vessel formation in the endometrium [14]. EPO stimulates proliferation and migration of human and bovine EC and also angiogenesis of the rat thoracic aorta [8, 15].

Many anemic patients with chronic renal failure receive treatment with recombinant human erythropoietin (rHuEPO) with beneficial results. The major side-effect of this treatment is the development of clinically significant hypertension [16–18]. Several factors have been implicated in rHuEPO-associated hypertension: an increase in hematocrit and red blood cell mass resulting in increased blood viscosity [19], a loss of hypoxic vasodilation [20], a direct vasoconstrictor effect [21], and an increase in calcium uptake by vascular smooth muscle cells [22, 23]. Nevertheless, the principal cause of EPO-induced hypertension is unknown. The angiogenic nature of EPO and the occurrence of EPOR on ECs suggest that EPO may directly stimulate ECs.

In humans, clinical events such as hypertension and thrombosis are often localized in specific vessels. This pattern, in part, may be due to the heterogeneity of EC themselves [reviewed in 24, 25]. Since the discovery of EC-dependent vasodilation by Furchgott and Zawadzki, ECs have been recognized as an important functional unit. Upon induction by vasoactive agents, such as acetylcholine and bradykinin, ECs secrete short-lived relaxing factor(s) causing relaxation of underlying smooth muscle cells [reviewed in 26, 27]. One endothelium-derived relaxing factor is nitric oxide (NO) [28]. EC-dependent vaso-

**Key words:** erythropoietin, receptor, endothelin, nitric oxide, anemia, chronic renal failure.

Received for publication August 5, 1999

and in revised form October 20, 1999

Accepted for publication December 2, 1999

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constriction has also been observed in response to various physiological stimuli such as thrombin [29], hypoxia [30], and mechanical stretch [31]. Endothelin-1 (ET-1) is the most potent vasoconstrictor peptide produced by ECs [32, 33]. Vascular smooth muscle cells and glomerular mesangial cells respond to ET-1 by cellular contraction and proliferation [34, 35]. To date, very little information is available on the effect of EPO on endothelium from different vascular sites.

In this study, we obtained primary cultures of EC prepared from various sites of human vasculature and examined the levels of EPO binding to the cells and the effect of EPO on EPOR, ET-1, and NO synthase (NOS) mRNA expression by quantitative reverse transcription-polymerase chain reaction (RT-PCR). We also measured the effect of EPO on ET-1 release into the culture medium and endothelial NOS enzyme activities. We found that EPORs are present on all ECs tested and that EPOR, ET-1, and NOS mRNA levels are unaffected by treatment with EPO. However, the extended exposure of vascular ECs to EPO at super-physiological concentrations induced NOS activity, which may contribute to the regulation of blood pressure.

## METHODS

### Cells and cell culture

Primary cultures of human ECs, prepared from umbilical vein (HUVEC), coronary artery (HCAEC), pulmonary artery (HPAEC), and dermis (HDEC) were purchased from Clonetics (San Diego, CA, USA). Cells were cultured at 37°C in modified MCDB 131 medium (Clonetics) supplemented with 5% fetal bovine serum (FBS; Hyclone Laboratories, Logan, UT, USA) in an air-5% CO<sub>2</sub> atmosphere at constant humidity. Cells were maintained in continuous culture and used within three to five passages.

### <sup>125</sup>I-EPO binding reactions

For binding studies (3-[<sup>125</sup>I]iodotyrosyl)rHuEPO (Amersham Corp., Arlington Heights, IL, USA) was used. The experimental procedures for the binding of <sup>125</sup>I-EPO to ECs have been reported elsewhere [8]. Briefly, confluent monolayers of ECs (1 × 10<sup>6</sup> cells/well) were washed with binding medium (0.05 mol/L phosphate buffer, pH 7.4, containing 150 mmol/L NaCl, 68 mmol/L CaCl<sub>2</sub>, 50 mmol/L MgCl<sub>2</sub>, and 1 mg/mL human serum albumin) and incubated at 22°C with 1.0 mL binding buffer containing 60,000 counts per min (cpm) of <sup>125</sup>I-EPO (Sp. Act. 300 to 900 Ci/mmol). At the end of specified time of incubation, wells were gently washed three times with prewarmed phosphate-buffered saline (PBS) containing 1% FBS, and the cells were then solubilized in lysis buffer [20 mmol/L HEPES, pH 7.4, 1% Triton X-100, 10% (vol/vol) glycerol, and 0.1 mg/mL of bovine serum albumin]. The radioactivity of the lysates was mea-

sured in a gamma counter. Nonspecific binding was assessed in the presence of a 100-fold molar excess of unlabeled EPO added at the start of incubation and subtracted from the total to calculate net specific binding.

### Preparation of RNA

Approximately 1 to 5 × 10<sup>7</sup> cells, either treated with rHuEPO (4 U/mL) for four hours or untreated, were washed twice with PBS, and RNA was extracted with TRIzol reagent as recommended by the supplier (Life Technologies, Grand Island, NY, USA). mRNA was adsorbed onto oligo(dT)-cellulose columns (Qiagen, Valencia, CA, USA). The total amount and concentration of mRNA were determined spectrophotometrically.

### Reverse transcription of RNA and synthesis of cDNA

An aliquot of mRNA (0.5 to 1 µg) was incubated at 42°C for one hour in 20 µL of 10 mmol/L Tris-HCl, pH 8.8, containing 50 mmol/L KCl, 0.1% Triton X-100, 5 mmol/L MgCl<sub>2</sub>, 1 mmol/L each dNTPs, 20 units of RNasin, and 0.5 µg oligo dT<sub>15</sub> primer. The cDNA was synthesized by primer extension using 15 units of avian myeloblastosis virus (AMV) reverse transcriptase per µg of RNA (Promega, Madison, WI, USA).

### Preparation of internal standard DNA and amplification of cDNA

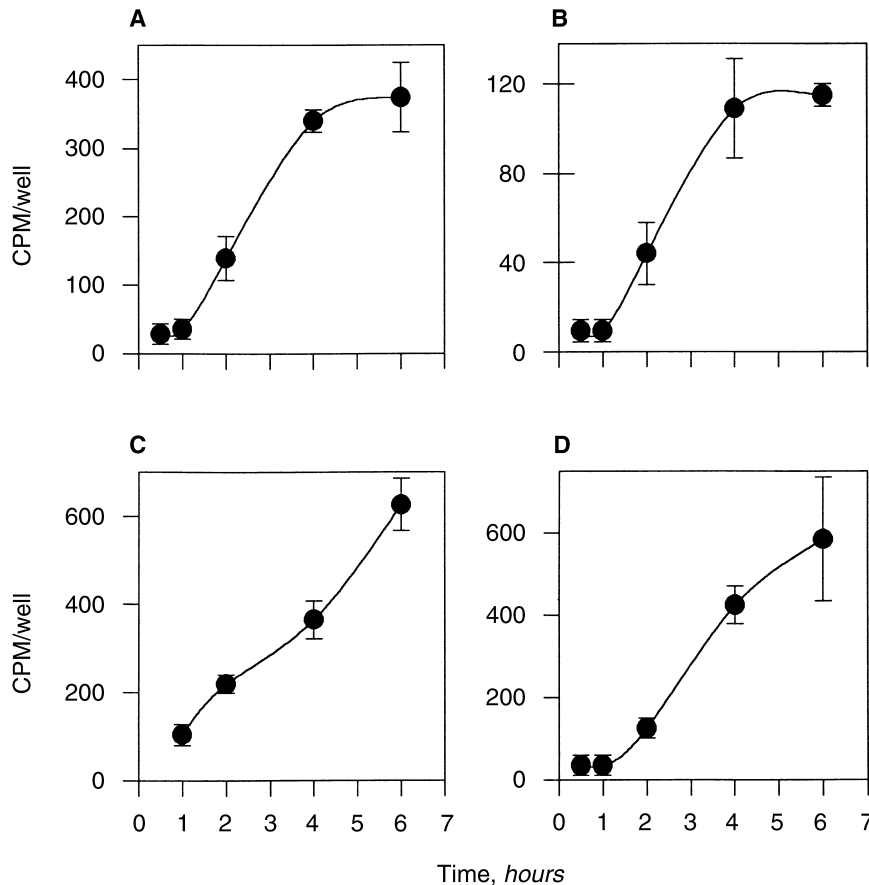
An internal standard was constructed by PCR amplification of genomic DNA using the following primer sets: (1) 5'-GCACCGAGTGTGTGCTGAGC-3' and 5'-GGT CAGCAGCACCAGGATGA-3' for hEPOR [36], (2) 5'-CCGTATGGACTTGGAAGCCC-3' and 5'-CTGG TTTGTCTTAGGTGTTCC-3' for ET-1 [37], and (3) 5'-CCTCCCCCGGCTGGGTGCG-3' and 5'-GCAC CTCCAGAAGCGTGCG-3' for NOS [38]. The same primer sets were used for cDNA amplification.

### Quantitative polymerase chain reaction

To quantitate gene-specific mRNA, multiple reaction mixtures were prepared with known amounts of cDNA. Serial dilutions of internal standard cDNA and [<sup>32</sup>P]CTP were added to each tube and coamplified. The products were analyzed by gel electrophoresis. The bands representing gene-specific cDNA and the standard DNA fragment were recovered and counted, and the results were plotted as cpm versus the concentration of external standard.

### Endothelin-1 assay

The medium from cultured ECs, either exposed for four hours to rHuEPO (4 U/mL) or untreated, was harvested and centrifuged at 500 × g at 4°C for 10 minutes. The recovered supernatants were then stored at -70°C. ET-1 determinations were performed using an ET-1 radioimmunoassay kit (Peninsula Lab., Belmont, CA, USA).



**Fig. 1. Kinetics of <sup>125</sup>I-recombinant human erythropoietin (rHuEPO) binding to endothelial cells.** Monolayers of human endothelial cells prepared from (A) umbilical vein (HUVEC), (B) coronary artery (HCAEC), (C) dermis (HDEC), and (D) pulmonary artery (HPAEC) were incubated with 60,000 cpm of <sup>125</sup>I-EPO (Sp. act. 166 Ci/mmol) in 1.0 mL phosphate buffered saline (PBS) containing CaCl<sub>2</sub> and MgCl<sub>2</sub> for increasing time intervals at 22°C. At the end of each time point, cells were washed, and the specific radioactivity associated with each well was determined as described in the **Methods** section. The results from three independent experiments are shown. Error bars represent standard errors (N = 3).

### Nitric oxide synthase assay

Confluent monolayers of ECs, either treated with rHuEPO (4 U/mL) daily for six days or untreated, were washed twice with PBS, harvested, and homogenized. The homogenate was centrifuged, and the supernatant protein concentration was adjusted to 1 mg/mL. The NOS activity was measured using a NOSdetect assay kit according to the supplier's specification (Stratagene, La Jolla, CA, USA).

## RESULTS

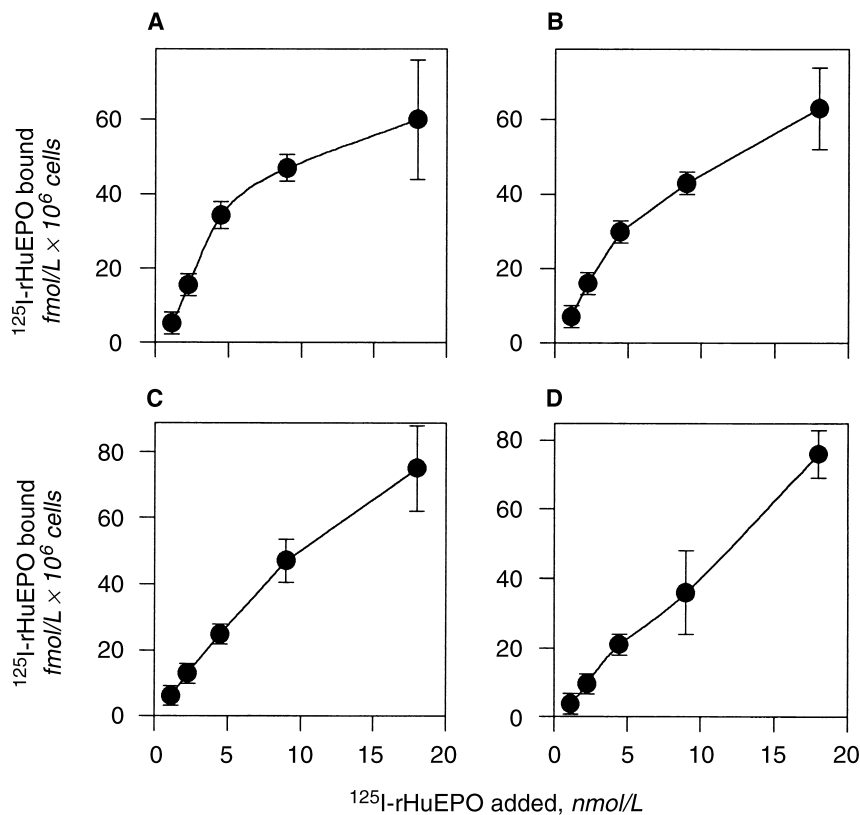
### <sup>125</sup>I-EPO binding study

To compare binding of <sup>125</sup>I-EPO to EC from various vascular sites, cells were exposed to <sup>125</sup>I-EPO with or without a 100-fold excess of unlabeled EPO for increasing time intervals. The time dependence of binding of <sup>125</sup>I-rHuEPO to ECs from various sources is shown in Figure 1. All ECs readily bound <sup>125</sup>I-EPO, and the specific binding increased with time, reaching a plateau in about four hours for HUVEC and HCAEC. The plateauing time was slightly longer (approximately 6 h) for HDEC and HPAEC. The fluctuation of increase binding in various ECs may correspond to the differences in the inter-

nalization and degradation of EPO as the binding assays were performed at 22°C. The concentration dependence of EPO binding was measured over the range from 1 to 18 nmol/L (Fig. 2). Specific binding was found to be saturable. The mean number of EPOR per cell was calculated by Scatchard plot analysis (Fig. 3). The calculated number of binding sites was 44,668 for HUVEC, 49,845 for HCAEC, 56,106 for HDEC, and 58,093 for HPAEC, while the dissociation constants ( $K_d$ ) were  $5.6 \times 10^{-9}$  mol/L for HUVEC,  $7.7 \times 10^{-9}$  mol/L for HCAEC,  $1.10 \times 10^{-8}$  mol/L for HDEC, and  $1.50 \times 10^{-8}$  mol/L for HPAEC. Exposure of cells for two days to EPO resulted in no significant change in the number of receptor sites (data not shown).

### Quantitation of mRNA expression

To confirm that EPO binding correlated with EPOR gene expression, ECs were cultured with or without EPO for four hours, and EPOR mRNA was measured by quantitative RT-PCR. The results are shown in Figure 4. The expected 197 bp cDNA fragment and 285 bp internal standard DNA were amplified from ECs from all sources. Further cloning and sequencing confirmed that the 197 bp fragment represented hEPOR mRNA.



**Fig. 2.** Dose response of  $^{125}\text{I}$ -EPO binding to endothelial cells. Confluent monolayers of (A) HUVEC, (B) HCAEC, (C) HDEC, and (D) HPAEC were washed and incubated with increasing amounts of  $^{125}\text{I}$ -EPO (1 to 18 pmol) for four hours at  $22^\circ\text{C}$ . Monolayers were washed, and the specific radioactivity associated with each well was determined as described in the **Methods** section. The results from three independent experiments are shown. Error bars represent standard errors ( $N = 3$ ).

The amount of hEPOR-specific cDNA/ $\mu\text{g}$  of mRNA determined in duplicate by quantitative PCR was  $5.1 \times 10^{-4}$  ng for HUVEC,  $11.0 \times 10^{-4}$  ng for HPAEC,  $4.0 \times 10^{-4}$  ng for HCAEC, and  $64.0 \times 10^{-4}$  for HDEC. With the exception of HCAEC, treatment of these cells with EPO for up to four hours had little or no effect on EPOR mRNA. In contrast, a twofold increase of EPOR mRNA was measured in EPO-treated HCAEC culture (Table 1). Treatment of HUVEC cells with EPO for 1, 2, 4, 6, and 24 hours had little or no effect on EPOR mRNA (data not shown).

#### Effects of rHuEPO on ET-1 gene expression and ET-1 secretion

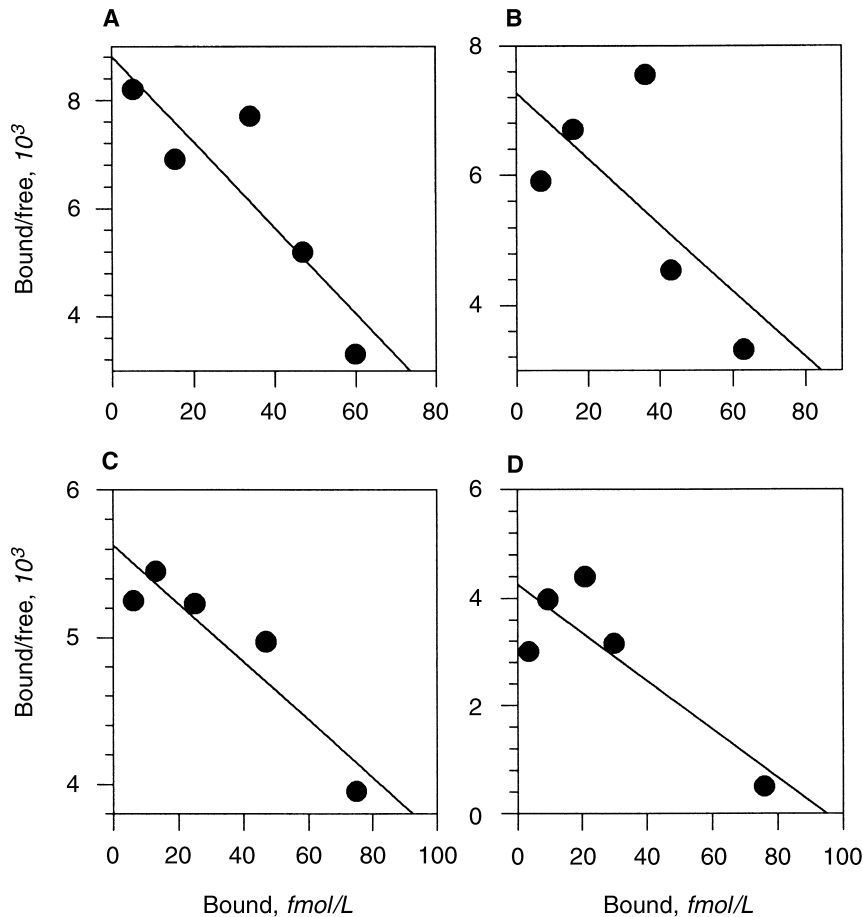
A direct link between ET-1 and EPO has been established from the findings that HUVECs have EPOR [8, 9] and that rHuEPO increases ET-1 release by bovine pulmonary artery ECs [39]. We therefore examined the effect of EPO on ET-1 expression by cultured HUVEC, HCAEC, HPAEC, and HDEC. As shown in Figure 5, a 290 bp ET-1-specific partial cDNA fragment and a 450 bp internal standard DNA were amplified by ET-1 primer sets and RT-PCR. The 290 bp fragment was cloned, and its identity confirmed by sequencing. The quantitative values are shown in Table 2. The amount of ET-1-specific cDNA/ $\mu\text{g}$  of mRNA in duplicate quantitative PCR was  $2.8 \times 10^{-2}$  ng for HUVEC,  $3.8 \times 10^{-2}$

ng for HPAEC,  $2.6 \times 10^{-2}$  ng for HCAEC, and  $0.34 \times 10^{-2}$  ng for HDEC. These values were unchanged in HCAEC, HPAEC, and HDEC cultures exposed to rHuEPO. However, the level of ET-1 transcript rose twofold in EPO-treated HUVEC cultures. Treatment of HUVEC cells with EPO for 1, 2, 4, 6, and 24 hours had no significant effect on ET-1 mRNA (data not shown).

Table 3 shows ET-1 release by cultured HUVEC, HCAEC, HPAEC, and HDEC during a four-hour treatment with 4 U/mL rHuEPO. All cells released ET-1 into the media. About 35 to 48 pg of ET-1 were released by  $2 \times 10^5$  cells. These values also did not change with EPO treatment, consistent with the results shown in Table 2.

#### Effect of rHuEPO on NOS activity

Nitric oxide is thought to play a central role in vascular homeostasis. NO, a heterodiatomic free radical product, is generated through the oxidation of L-arginine to L-citrulline by NOS. NO has been demonstrated to inhibit thrombosis, cytokine-induced vascular cell adhesion molecule-1 (VCAM) expression, leukocyte adhesion to endothelium, and smooth muscle proliferation and migration [40–42]. To investigate the effect of EPO on NOS activity, we measured NOS mRNA levels by RT-PCR. The results are shown in Figure 6. Partial cDNA fragments (approximately 230 bp) and standard DNA (approximately 500 bp) were isolated, cloned, and char-



**Fig. 3. Scatchard analysis of  $^{125}\text{I}$ -EPO binding to endothelial cells.** Plots represent Scatchard analyses of binding data shown in Figure 2. The data suggest the presence of a single class receptor with different affinities in different cell types. For (A) HUVEC,  $K_d = 5.6 \times 10^{-9}$  mol/L and 44,668 receptor sites per cell; (B) HCAEC,  $K_d = 7.7 \times 10^{-9}$  mol/L and 49,845 receptor sites per cell; (C) HDEC,  $K_d = 1.10 \times 10^{-8}$  mol/L and 56,106 receptor sites per cell; (D) HPAEC,  $K_d = 1.5 \times 10^{-8}$  mol/L and 58,093 receptor sites per cell.

acterized by sequencing. The quantitative values are shown in Table 4. The NOS-specific mRNA in ECs varied from  $1.0 \times 10^{-3}$  to  $1.8 \times 10^{-3}$  ng/ $\mu\text{g}$  of mRNA. Exposure of the cells to rHuEPO had little effect. However, the presence of EPO for 24 hours induced a three-fold increase in NOS-specific mRNA ( $5.4 \times 10^{-3}$  ng cDNA/ $\mu\text{g}$  of mRNA) compared with control cultures of HUVEC ( $1.8 \times 10^{-3}$  ng cDNA/ $\mu\text{g}$  of mRNA). The increase values were maintained for up to six days upon daily removal and replenishment with fresh EPO-supplemented medium. Short-term exposure for one, two, four, and six hours to EPO, HUVEC exhibited no effect on NOS-specific mRNA (data not shown).

To confirm that the increased NOS-specific mRNA resulted in increased NOS activity, we determined the effect of EPO on NOS activity. As shown in Table 5, long-term EPO treatment resulted in a twofold to fourfold increase in NOS activity in all ECs.

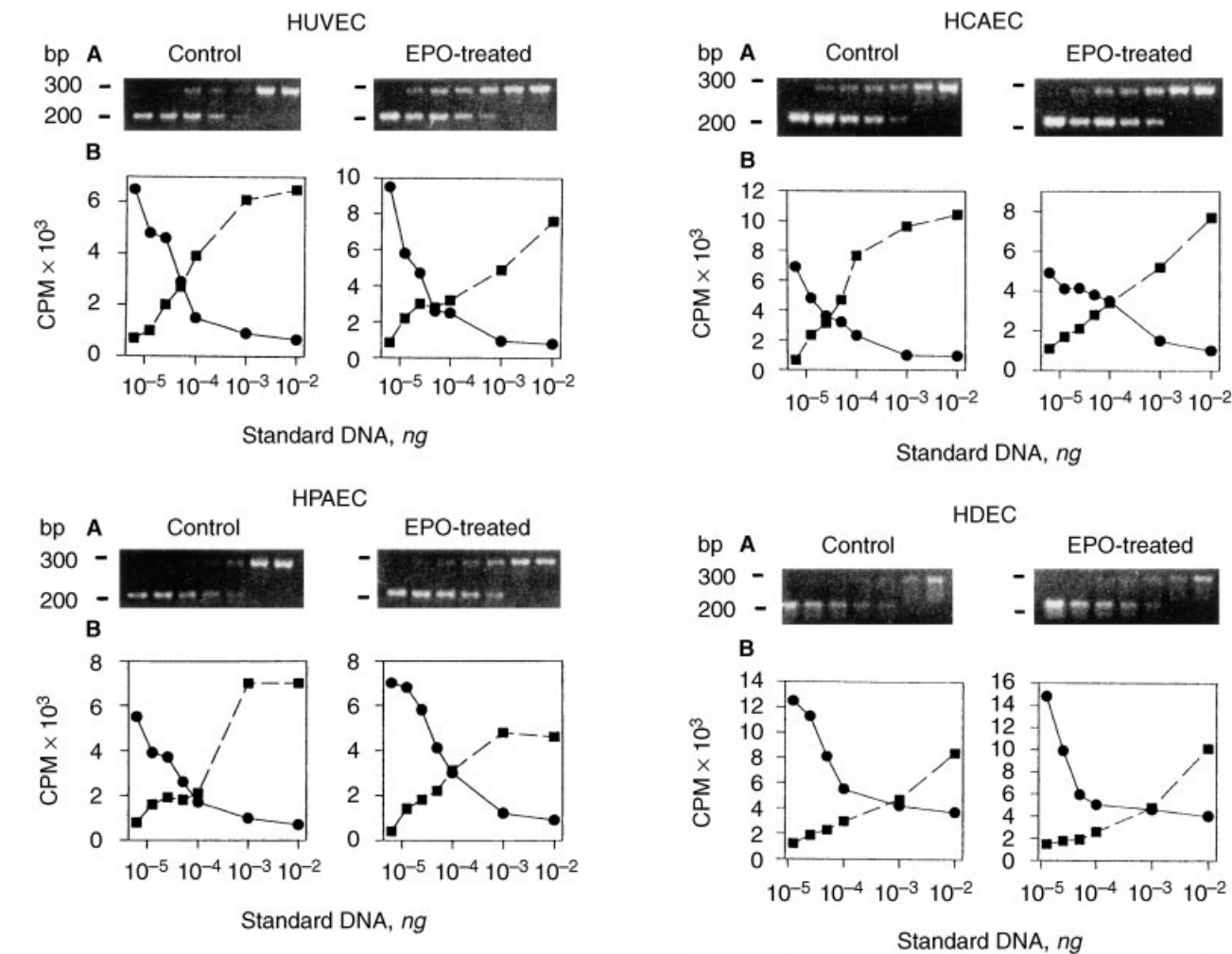
## DISCUSSION

The present study demonstrates that ECs prepared from various regions of the vascular tree bind EPO in a dose-dependent manner. The scattering of data points

at a low ligand concentration in Scatchard analysis indicates the possibility of more than one binding site. Our finding of a large number of relatively low-affinity binding sites on all EC types tested (Fig. 3) is consistent with the findings of Anagnostou et al [8]. This is in contrast with earlier studies that measured the binding of EPO to its receptors and showed that erythroid progenitor cells have only a small number of cell surface receptors [43–45]. The low number of receptors per cell for growth factors is common in hematopoietic cells [46].

The ligand affinity of EPOR expressed on EC is much lower than that on erythroid cells; the  $K_d$  for EPO binding to EPOR on EC varies from approximately 5 to 15 nmol/L [8, 10], while the  $K_d$  for the high-affinity receptor on erythroid cells is approximately 50 pmol/L [47]. The reason for the expression of only the low-affinity receptor on ECs is unclear. Expression of only the low-affinity receptor might be crucial to prevent unregulated angiogenesis by endogenous EPO. The presence of EPOR on ECs may originate from the close relationship between vasculogenesis and hematopoiesis in early development [48] and between angiogenesis and hematopoiesis later in development [49].





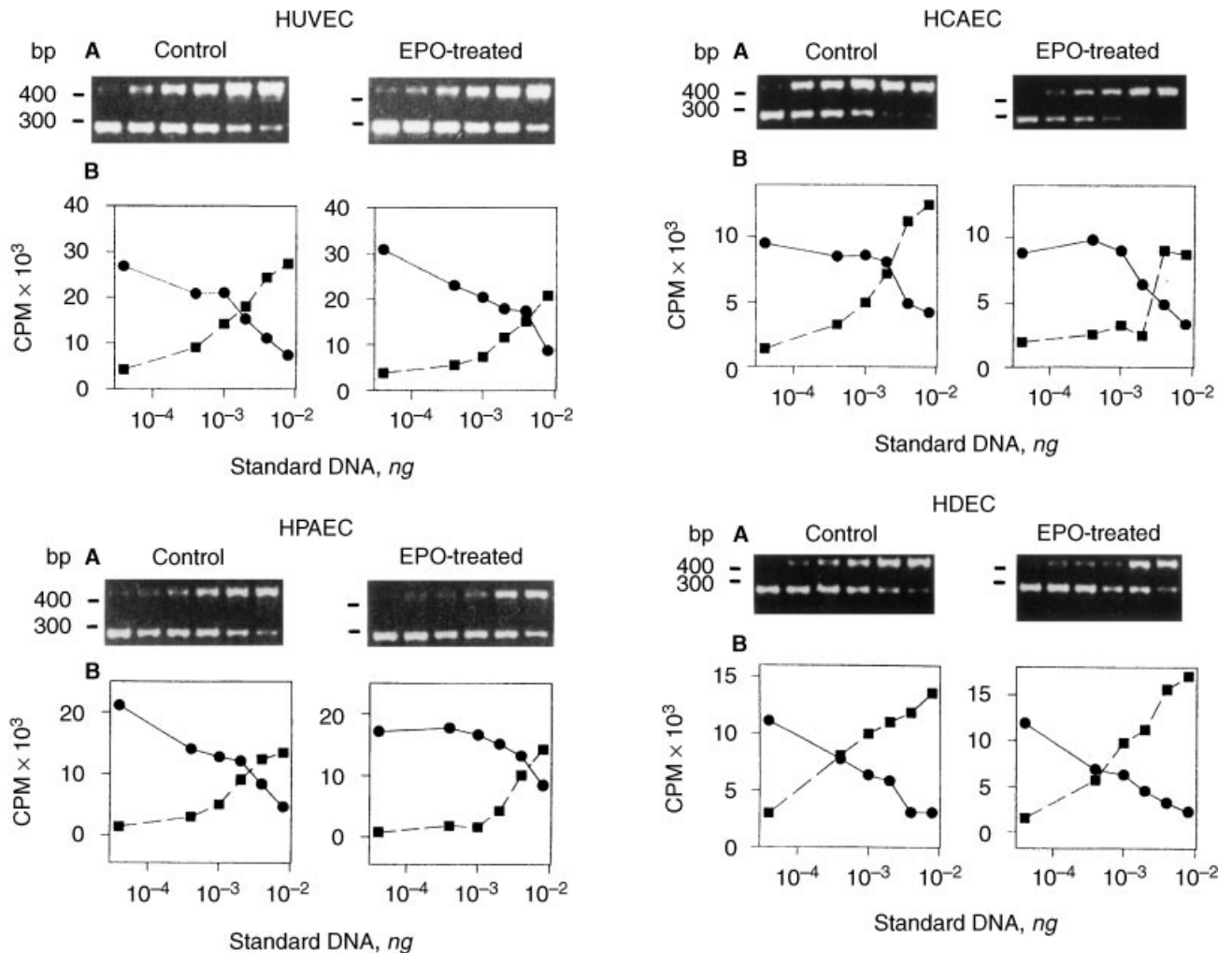
**Fig. 4. Polymerase chain reaction (PCR) analysis of human erythropoietin receptor (hEPOR) transcripts in control and EPO-treated HUVEC, HPAEC, HDEC, and HCAEC.** Monolayers of HUVEC, HCAEC, HDEC, and HPAEC were incubated with EPO (4 U/mL) for four hours or were untreated, and mRNA was prepared as described in the **Methods** section. cDNA was reverse transcribed from mRNA isolated from respective endothelial cells. Multiple reactions for hEPOR-specific PCR amplification of cDNA were carried out with hEPOR primers. Prior to amplification, increasing amounts of standard DNA and [ $\alpha$ -<sup>32</sup>P]dCTP were added to each reaction. (A) The amplification products were resolved by agarose gel electrophoresis. Left panel show control gels and right panels show EPO-treated gels (upper band 285 bp standard DNA and lower band 197 bp hEPOR cDNA; bp, size markers). (B) For quantitation, the hEPOR and the standard DNA bands were cut, the radioactivity present in each band was determined and plotted against the amount of standard added to each reaction mixture. Symbols are: (●) standard DNA; (■) hEPOR-specific PCR product. The amount of standard corresponding to the point at which the two PCR products are equal is an indication of the amount of hEPOR cDNA present.

ECs	Control	EPO-treated
	ng cDNA/ $\mu$ g of mRNA	
HUVEC	$5.1 \times 10^{-4}$	$5.5 \times 10^{-4}$
HPAEC	$11 \times 10^{-4}$	$9.2 \times 10^{-4}$
HCAEC	$4.0 \times 10^{-4}$	$9.0 \times 10^{-4}$
HDEC	$64.0 \times 10^{-4}$	$88.0 \times 10^{-4}$

Abbreviations are: ECs, endothelial cells; EPO, erythropoietin; HUVEC, human umbilical vein endothelial cells; HPAEC, human pulmonary artery endothelial cells; HCAEC, human coronary artery endothelial cells; HDEC, human dermis endothelial cells. The values of human EPO receptor-specific mRNA were obtained from Figure 4B. Each sample was assayed three times from two separate experiments.

Compared with OCI-M1 or other human erythroid cell lines, HUVECs have many more EPOR and a lower expression of EPOR mRNA [9]. In our study, the mean number of EPOR per cell varied from 44,668 to 58,093, but the quantity of EPOR mRNA was less than the amount reported for OCI-M1 cells. The significance of this discrepancy is not clear. In erythroid cells, less than 5% of the newly synthesized EPORs are found on the cell surface, and the rest are degraded in the endoplasmic reticulum [50, 51].

Quantitative determination of mRNA expression failed to detect any increase in EPOR gene expression in EPO-treated HUVEC, HPAEC, and HDEC cultures. In contrast, a twofold increase of EPOR mRNA was



**Fig. 5. Polymerase chain reaction (PCR) analysis of endothelin-1 (ET-1) transcripts in untreated and EPO-treated HUVEC, HPAEC, HDEC, and HCAEC.** Monolayers of HUVEC, HCAEC, HDEC, and HPAEC cells were either incubated with EPO (4 U/mL) for four hours or were untreated, and mRNA was prepared as described in the **Methods** section. Reverse transcription of total RNA isolated from respective endothelial cells with ET-1-specific primers and multiple reactions for PCR amplification of cDNA were carried out with ET-1 primers and standard DNA as described in Figure 4. (A) The amplification products were resolved by agarose gel electrophoresis. Left panels show controls and right panels show EPO treated (upper band 450 bp standard DNA and lower band 290 bp ET-1 cDNA; bp, size markers). (B) Quantitation was achieved as described in Figure 4. ET-1 cDNA and the standard DNA bands were cut, the radioactivity present in each band was determined and plotted against the amount of standard added to each reaction mixture. Symbols are: (●) standard DNA; (■) ET-1-specific PCR product to each reaction mixture.

**Table 2.** Quantitation of endothelin-1 (ET-1)-specific mRNA expression

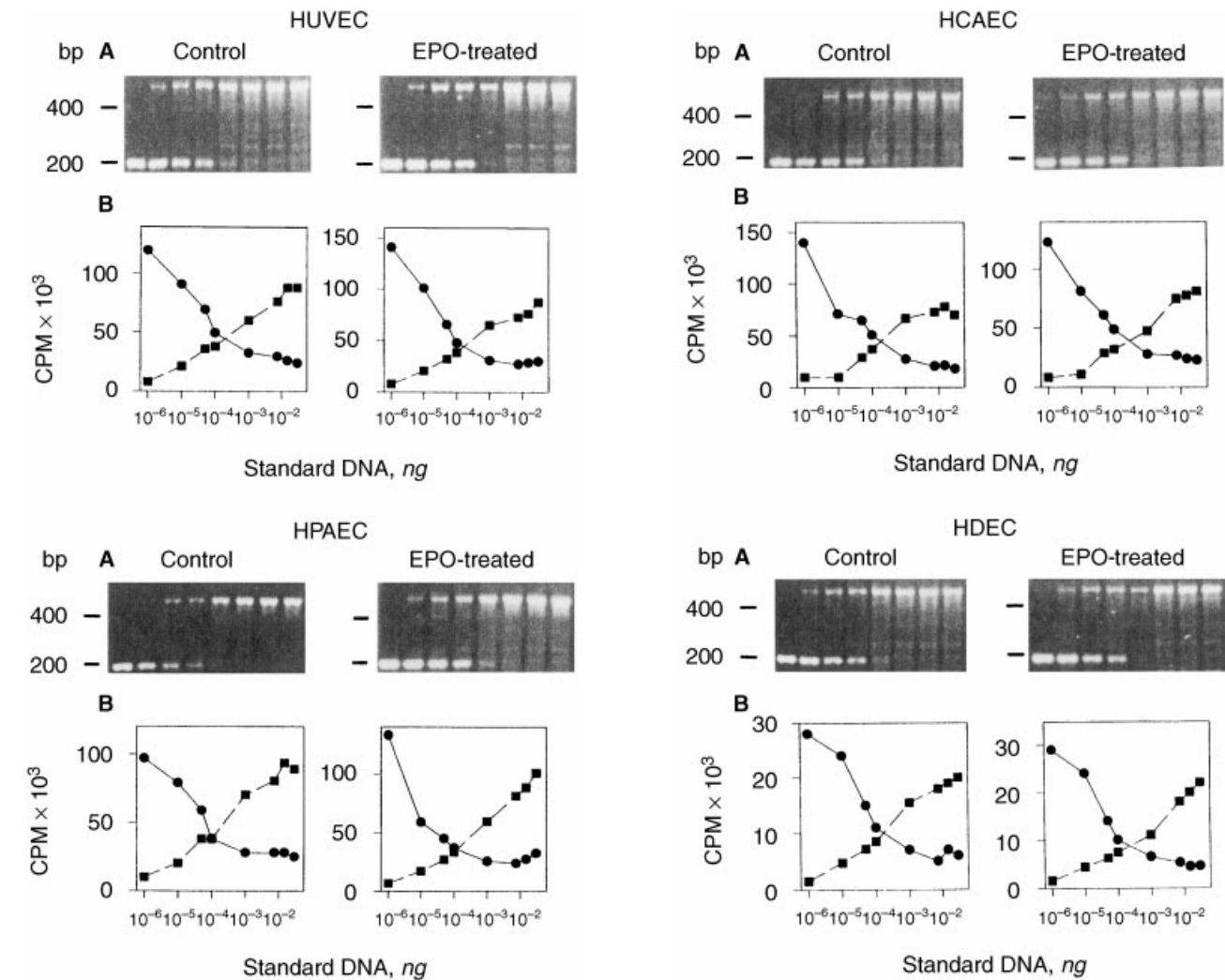
	Control	EPO-treated
ECs	<i>ng cDNA/μg of mRNA</i>	
HUVEC	$2.8 \times 10^{-2}$	$6.2 \times 10^{-2}$
HPAEC	$3.8 \times 10^{-2}$	$4.7 \times 10^{-2}$
HCAEC	$2.6 \times 10^{-2}$	$2.7 \times 10^{-2}$
HDEC	$0.34 \times 10^{-2}$	$0.43 \times 10^{-2}$

The values of human ET-1-specific mRNA were derived from Figure 5B. Each sample was assayed three times from two separate experiments. Abbreviations are in Table 1.

**Table 3.** Effect of EPO on ET-1 production by endothelial cells from various sources

	Control	EPO-treated
ECs	<i>Endothelin-1 pg/2 × 10<sup>5</sup> cells</i>	
HUVEC	35	37
HPAEC	39	36
HCAEC	48	50
HDEC	45	47

HUVEC, HPAEC, HCAEC, and HDEC were incubated in serum-free endothelial cell basal media (EBM) with 4 U/mL rHuEPO for 4 hours. Control cells were exposed to rHuEPO vehicle alone. Results represent the mean of two experiments performed in duplicate. Abbreviations are in Table 1.



**Fig. 6.** Polymerase chain reaction (PCR) analysis of nitric oxide synthase (NOS) transcripts in EPO-treated and nontreated HUVEC, HPAEC, HDEC, and HCAEC. Monolayers of HUVEC, HCAEC, HDEC, and HPAEC were either incubated with EPO (4 U/mL) for four hours or were untreated, and mRNA was prepared as described in the **Methods** section. Reverse transcription of total RNA isolated from respective endothelial cells with NOS-specific primers and multiple reactions for PCR amplification of cDNA were carried out with NOS primers and standard DNA, as described in Figure 5. (A) The amplification products were analyzed by agarose gel electrophoresis. Left panels show controls and right panels show EPO treated (upper 500 bp band standard DNA and lower 200 bp NOS cDNA band; bp, size markers). (B) For quantitation, the NOS and the standard bands were cut, the radioactivity present in each band was determined and plotted against the amount of standard added to each reaction mixture. Symbols are: (●) standard DNA; (■) NOS-specific PCR product.

**Table 4.** Quantitation of nitric oxide synthase (NOS)-specific mRNA expression

	Control	EPO-treated
ECs	ng cDNA/ $\mu$ g of mRNA	
HUVEC	$1.8 \times 10^{-3}$	$1.6 \times 10^{-3}$
HPAEC	$1.0 \times 10^{-3}$	$0.8 \times 10^{-3}$
HCAEC	$1.8 \times 10^{-3}$	$1.9 \times 10^{-3}$
HDEC	$1.2 \times 10^{-3}$	$1.5 \times 10^{-3}$

The values of human NOS-specific mRNA were obtained from Figure 6B. Each sample was assayed three times from two separate experiments. Abbreviations are in Table 1.

**Table 5.** Effect of EPO on NOS activity by endothelial cells from various sources

	Control	EPO-treated	Change(-fold)
ECs	Nitric oxide synthase $\text{fmol}^3\text{H-citrulline/min/mg protein}$		
HUVEC	30	78	2.6
HPAEC	14	63	4.5
HDEC	30	110	3.6

HUVEC, HPAEC, and HDEC were incubated in complete media (EGM-2 MV) with 4 U/mL rHuEPO. The medium was changed daily for up to 6 days. Control cells were exposed to complete media alone. The cell extracts were prepared and NOS activity was measured as described in the **Methods** section. Results represent the mean of two experiments performed in duplicate. Abbreviations are in Table 1.



found in EPO-treated HCAEC culture (Table 1). Similar determinations of ET-1 transcripts revealed no increase in ET-1 mRNA in EPO-treated HPAEC, HCAEC, and HDEC cultures, whereas a twofold increase of ET-1 mRNA was found in EPO-treated HUVEC (Table 2). The reasons for these differences are not clear. It is possible that the untreated HCAEC and HUVEC cultures contained fewer ECs, and thus, the technique can be limited by its inability to distinguish heterogeneous cell populations.

Normally, ECs contribute to the regulation of blood pressure and blood flow by releasing vasodilators such as NO and prostacyclin (PGI<sub>2</sub>), as well as vasoconstrictors including ET-1 and platelet activating factor (PAF) [reviewed in 52]. ECs synthesize ET-1 [53]. ET-1 is not stored in granules but is transcribed after stimulation by hypoxia, shear stress, or ischemia [54]. rHuEPO has been shown to have a direct stimulatory effect on ET-1 release from cultured bovine PAEC [39]. This stimulatory effect was time dependent, peaked at 4 hours, and reached a plateau at 12 hours. In contrast, we saw no effect on ET-1 release in response to EPO in any of the four human EC types in culture. This may be due to differences in culture conditions.

Endothelial cells produce NO, a free radical generated through the oxidation of L-arginine to L-citrulline by NO synthase [55]. One isoform, eNOS, is constitutively active in EC and can be further stimulated by thrombin, bradykinin, or shear stress [56]. Our study demonstrates that short-term treatment of ECs in culture has little or no effect on NOS transcript. However, EC cultures exposed to EPO for several days increased their production of NO. Although we did not measure NO release into the medium and it is impossible to directly extrapolate these results to the in vivo situation, the increased production of NO may contribute to the regulation of blood pressure.

Heidenreich, Rahn, and Zidek [21] and Carlini et al [39] have shown that rHuEPO has a direct vasoconstrictive effect. In addition, in vivo studies of Buemi, Allegra, and Frisina [57] and the results of the present studies (Table 5) are consistent with a vasodilatation effects of EPO. EPO may act by modifying the balance of vasoconstrictive and vasodilatory effects to favor vasoconstriction in the setting of renal failure. Why hypertension is only seen in this patient group remains unresolved.

## ACKNOWLEDGMENTS

We wish to thank Mr. Tellervo Huima and Ms. Yelena Oksov for the preparation of the illustrations.

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